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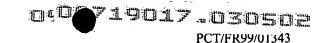
## SEQUENCE LISTING

<110> Pierrard, Jerome Guitton, Carole Favre-Bulle, Olivier <120> Industrial Method For Producing Heterologous Proteins In E. coli And Strains Useful For Said Method <130> 5500\*54 <140> US 09/719,017 <141> 1999-06-08 <150> PCT/FR99/01343 <151> 1999-06-08 <150> FR 98/07,474 <151> 1998-06-10 <160> 6 <170> PatentIn Ver. 2.1 <210> 1 <211> 121 <212> DNA <213> Escherichia coli <400> 1 gaatteeetg ttgacaatta atcategaac tagttaacta gtacgeaget tggetgeagg 60 tcgacctgca gccaagcttg ggcatacatt caatcaattg ttatctaagg aaatacttac 120 121 <210> 2 <211> 1793 <212> DNA <213> Alcaligenes faecalis <220> <221> CDS <222> (123)..(1190) <400> 2 qaatteeetq ttqacaatta atcategaac tagttaacta gtacgeaget tggctgcagg 60 tcgacctgca gccaagcttg ggcatacatt caatcaattg ttatctaagg aaatacttac 120 at atg cag aca aga aaa atc gtc cgg gca gcc gcc gta cag gcc gcc Met Gln Thr Arg Lys Ile Val Arg Ala Ala Ala Val Gln Ala Ala 1 tct ccc aac tac gat ctg gca acg ggt gtt gat aaa acc att gag ctg

Ser Pro Asn Tyr Asp Leu Ala Thr Gly Val Asp Lys Thr Ile Glu Leu

25

20



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ggc ctg atc att Gly Leu Ile Ile 275			Glu Glu	Ile Ala P		983
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ctg gta ctg gac Leu Val Leu Asp 305	Leu Gly H					1079
aaa agc gtg atc Lys Ser Val Ile 320						1127
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Val Gln Glu Pro	Ser					
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Val Gln Glu Pro 355 tcttgattcc ttct	Ser gcgtcc cgg gacccc atg	atccact ag ccgaact ca	ttctagag gaagtgaa	tcgacctgc	a ggcatgcaag c gccgatggta	1290 1350
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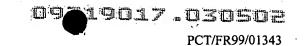
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Trp Leu Pro Gly Tyr Pro Phe His Val Trp Leu Gly Ala Pro Ala Trp 50 55 60

Ser Leu Lys Tyr Ser Ala Arg Tyr Tyr Ala Asn Ser Leu Ser Leu Asp 65 70 75 80

Ser Ala Glu Phe Gln Arg Ile Ala Gln Ala Ala Arg Thr Leu Gly Ile 85 90 95

Phe Ile Ala Leu Gly Tyr Ser Glu Arg Ser Gly Gly Ser Leu Tyr Leu 100 105 110

Gly Gln Cys Leu Ile Asp Asp Lys Gly Gln Met Leu Trp Ser Arg Arg 115 120 125

Lys Leu Lys Pro Thr His Val Glu Arg Thr Val Phe Gly Glu Gly Tyr 130 135 140

Ala Arg Asp Leu Ile Val Ser Asp Thr Glu Leu Gly Arg Val Gly Ala 145 150 155 160

Leu Cys Cys Trp Glu His Leu Ser Pro Leu Ser Lys Tyr Ala Leu Tyr 165 170 175

Ser Gln His Glu Ala Ile His Ile Ala Ala Trp Pro Ser Phe Ser Leu 180 185 190

Tyr Ser Glu Gln Ala His Ala Leu Ser Ala Lys Val Asn Met Ala Ala 195 200 205

Ser Gln Ile Tyr Ser Val Glu Gly Gln Cys Phe Thr Ile Ala Ala Ser 210 215 220

Ser Val Val Thr Gln Glu Thr Leu Asp Met Leu Glu Val Gly Glu His 225 230 235 240

Asn Ala Ser Leu Leu Lys Val Gly Gly Gly Ser Ser Met Ile Phe Ala 245 250 255

Pro Asp Gly Arg Thr Leu Ala Pro Tyr Leu Pro His Asp Ala Glu Gly 260 265 270

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